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=> s fusion (w)protein?
L1 118936 FUSION (W) PROTEIN?

=> s 11 (p)short(p)live?
L2 208 L1 (P) SHORT(P) LIVE?

=> s 12 (p)express?
L3 120 L2 (P) EXPRESS?

=> s 13 (p)select? (p)cell
L4 12 L3 (P) SELECT? (P) CELL

=> d his

(FILE 'HOME' ENTERED AT 12:17:06 ON 27 JAN 2003)

FILE 'MEDLINE, BIOSIS, CAPLUS, EMBASE' ENTERED AT 12:17:21 ON 27 JAN 2003
L1 118936 S FUSION (W) PROTEIN?
L2 208 S L1 (P) SHORT(P) LIVE?
L3 120 S L2 (P) EXPRESS?
L4 12 S L3 (P) SELECT? (P) CELL

=> duplicate remove 14
DUPLICATE PREFERENCE IS 'MEDLINE, BIOSIS, CAPLUS, EMBASE'
KEEP DUPLICATES FROM MORE THAN ONE FILE? Y/(N):n
PROCESSING COMPLETED FOR L4
L5 4 DUPLICATE REMOVE L4 (8 DUPLICATES REMOVED)

=> d his

(FILE 'HOME' ENTERED AT 12:17:06 ON 27 JAN 2003)

FILE 'MEDLINE, BIOSIS, CAPLUS, EMBASE' ENTERED AT 12:17:21 ON 27 JAN 2003
L1 118936 S FUSION (W) PROTEIN?
L2 208 S L1 (P) SHORT(P) LIVE?
L3 120 S L2 (P) EXPRESS?
L4 12 S L3 (P) SELECT? (P) CELL
L5 4 DUPLICATE REMOVE L4 (8 DUPLICATES REMOVED)

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L2 same express\$ 18

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side by side**Hit Count** Set Name
result set*DB=USPT; PLUR=YES; OP=OR*

<u>L3</u>	L2 same express\$	18	<u>L3</u>
<u>L2</u>	L1 same short same live\$	35	<u>L2</u>
<u>L1</u>	fusion near0 protein	14450	<u>L1</u>

END OF SEARCH HISTORY

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L1 same fusion same protein

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side by side**Hit Count** **Set Name**
result set*DB=USPT; PLUR=YES; OP=OR*

<u>L4</u>	L3 same (advantag\$ or useful\$)	0	<u>L4</u>
<u>L3</u>	L1 same fusion same protein	11	<u>L3</u>
<u>L2</u>	L1 same fusio same protein	0	<u>L2</u>
<u>L1</u>	pulse near0 chase near0 analysis	82	<u>L1</u>

END OF SEARCH HISTORY